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RAW SEQUENCE LISTING
ERROR REPORT

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Application Serial Number: 10/669,162 B
Source: FE016
Date Processed by STIC: 7/28/04

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

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Revised 05/17/04

RAW SEQUENCE LISTING

DATE: 07/28/2004

PATENT APPLICATION: US/10/669,162B

TIME: 15:47:40

Input Set : A:\228234_5.doc

Output Set: N:\CRF4\07282004\J669162B.raw

183 <212> TYPE: RNA
 184 <213> ORGANISM: Artificial Sequence
 186 <220> FEATURE:
 187 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
 188 synthetic construct
 190 <220> FEATURE:
 191 <221> NAME/KEY: misc_feature
 192 <222> LOCATION: 1, 7-9, 13, 14, 16, 18, 25, 26, 32, 33, 37, 39, 42, 43, 50,
 193 51
 194 <223> OTHER INFORMATION: n = g, a, c or u
 196 <220> FEATURE:
 197 <221> NAME/KEY: misc_feature
 198 <222> LOCATION: 38, 44
 199 <223> OTHER INFORMATION: r = a or g
 201 <220> FEATURE:
 202 <221> NAME/KEY: misc_feature
 203 <222> LOCATION: 17, 34
 204 <223> OTHER INFORMATION: y = c or u
 206 <400> SEQUENCE: 9
 > 207 ncuaaunnng agnngnynga gggannggcc cnnyganrnc cnrgcaach n SAME ERROR → 51
 365 <210> SEQ ID NO: 18
 366 <211> LENGTH: 22
 367 <212> TYPE: RNA
 368 <213> ORGANISM: Arabidopsis thaliana
 370 <220> FEATURE:
 371 <221> NAME/KEY: misc_feature
 372 <222> LOCATION: 9
 373 <223> OTHER INFORMATION: d = g, a or u
 375 <220> FEATURE:
 376 <221> NAME/KEY: misc_feature
 377 <222> LOCATION: 1-7, 10-16
 378 <223> OTHER INFORMATION: n = g, a, c or u
 380 <400> SEQUENCE: 18
 > 381 nnnnnngdn nnnnnncuga ga SAME ERROR → 22
 1043 <210> SEQ ID NO: 60
 1044 <211> LENGTH: 23
 1045 <212> TYPE: RNA
 1046 <213> ORGANISM: Bacillus subtilis
 1048 <220> FEATURE:
 1049 <221> NAME/KEY: misc_feature
 1050 <222> LOCATION: 12-131
 1051 <223> OTHER INFORMATION: n = g, a, c or u
 1053 <220> FEATURE:
 1054 <221> NAME/KEY: misc_feature
 1055 <222> LOCATION: 2, 11, 52, 53, 70, 92, 132
 1056 <223> OTHER INFORMATION: r = a or g
 1058 <220> FEATURE:
 1059 <221> NAME/KEY: misc_feature
 1060 <222> LOCATION: 3, 135

✓ Insert =

✓ Insert

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TIME: 15:47:40

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1061 <223> OTHER INFORMATION: w = a or u
 1063 <220> FEATURE:
 1064 <221> NAME/KEY: misc_feature
 1065 <222> LOCATION: 64, 72, 93, 119, 136
 1066 <223> OTHER INFORMATION: y = c or u
 1068 <400> SEQUENCE: 60 *Same error → 23*
 > 1069 rwagagggc rnnnnnnannn aqua
 1636 <210> SEQ ID NO: 98
 1637 <211> LENGTH: 51
 1638 <212> TYPE: RNA
 1639 <213> ORGANISM: Bacillus subtilis
 1641 <221> NAME/KEY: misc_feature
 1642 <222> LOCATION: 8, 13, 14, 26, 32, 33, 37, 41, 42, 50, 51, 54, 55, 63, 67
 1643 <223> OTHER INFORMATION: n = g, a, c or u
 1645 <220> FEATURE:
 1646 <221> NAME/KEY: misc_feature
 1647 <222> LOCATION: 18, 38, 44, 53, 68, 71, 72, 78, 79, 84, 87
 1648 <223> OTHER INFORMATION: r = a or g
 1650 <220> FEATURE:
 1651 <221> NAME/KEY: misc_feature
 1652 <222> LOCATION: 1, 17, 25, 34, 60, 74, 75
 1653 <223> OTHER INFORMATION: y = c or u
 > 1655 <400> 98
 > 1656 ycuuauacnag agunggyrga gggaynggcc cnnyganrcc nncrgcaacn n *same error → 51*
 7181 <210> SEQ ID NO: 379
 7182 <211> LENGTH: 30 *Found 31*
 7183 <212> TYPE: RNA
 7184 <213> ORGANISM: Artificial Sequence
 7186 <220> FEATURE:
 7187 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
 7188 synthetic construct
 7190 <220> FEATURE:
 7191 <221> NAME/KEY: misc_feature
 7192 <222> LOCATION: 3-4, 7-9, 12, 14-15, 21, 24, 28-30
 7193 <223> OTHER INFORMATION: n = g, a, c or u
 7195 <220> FEATURE:
 7196 <221> NAME/KEY: misc_feature
 7197 <222> LOCATION: 1, 10, 22, 27, 31
 7198 <223> OTHER INFORMATION: r = a or g
 7200 <400> SEQUENCE: 379
 > 7201 rannccnnnr cngnnccgc nrcngurnnn r 31 ✓
 7392 <210> SEQ ID NO: 388
 7393 <211> LENGTH: 19 *Found 18*
 7394 <212> TYPE: RNA
 7395 <213> ORGANISM: Artificial Sequence
 7397 <220> FEATURE:
 7398 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
 7399 synthetic construct
 7401 <220> FEATURE:

RAW SEQUENCE LISTING

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Input Set : A:\228234_5.doc

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7402 <221> NAME/KEY: misc_feature
 7403 <222> LOCATION: 1-3, 10-11, 14-17, 19
 7404 <223> OTHER INFORMATION: n = g, a, c or u
 7406 <220> FEATURE:
 7407 <221> NAME/KEY: misc_feature
 7408 <222> LOCATION: 18
 7409 <223> OTHER INFORMATION: r = a or g
 7411 <220> FEATURE:
 7412 <221> NAME/KEY: misc_feature
 7413 <222> LOCATION: 8
 7414 <223> OTHER INFORMATION: y = c or u
 7416 <400> SEQUENCE: 388
 7417 ~~nnnggcyann~~ ¹⁸ ~~nnnnnnnn~~ ¹⁹
 7498 <210> SEQ ID NO: 393
 7499 <211> LENGTH: 43 ^{Found 44}
 7500 <212> TYPE: RNA
 7501 <213> ORGANISM: Artificial Sequence
 7503 <220> FEATURE:
 7504 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
 7505 synthetic construct
 7507 <220> FEATURE:
 7508 <221> NAME/KEY: misc_feature
 7509 <222> LOCATION: 1-8, 14-20, 21-22, 32-43
 7510 <223> OTHER INFORMATION: n = g, a, c or u
 7512 <220> FEATURE:
 7513 <221> NAME/KEY: misc_feature
 7514 <222> LOCATION: 9-10, 29
 7515 <223> OTHER INFORMATION: r = a or g
 7517 <220> FEATURE:
 7518 <221> NAME/KEY: misc_feature
 7519 <222> LOCATION: 23, 31
 7520 <223> OTHER INFORMATION: y = c or u
 7522 <400> SEQUENCE: 393
 7523 ~~nnnnnnnnrrr. agggnnnnnnn nnyggcggarg ynnnnnnnnnnn nnn~~ ⁴³ ^{Found 44}
 7567 <210> SEQ ID NO: 396
 7568 <211> LENGTH: 27 ^{Found 17}
 7569 <212> TYPE: RNA
 7570 <213> ORGANISM: Artificial Sequence
 7572 <220> FEATURE:
 7573 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
 7574 synthetic construct
 7576 <220> FEATURE:
 7577 <221> NAME/KEY: misc_feature
 7578 <222> LOCATION: 1-11
 7579 <223> OTHER INFORMATION: n = g, a, c or u
 7581 <220> FEATURE:
 7582 <221> NAME/KEY: misc_feature
 7583 <222> LOCATION: 12
 7584 <223> OTHER INFORMATION: y = c or u

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/669,162B

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TIME: 15:47:41

Input Set : A:\228234 5.doc

Output Set: N:\CRF4\07282004\J669162B.raw

7586 <400> SEQUENCE: 396

--> 7587 ~~nyuguca~~ nyuguca

27-17

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/669,162B

DATE: 07/28/2004

TIME: 15:47:42

Input Set : A:\228234_5.doc

Output Set: N:\CRF4\07282004\J669162B.raw

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.:71 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:120
l:341 Repeated in SeqNo=3
.:109 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
l:341 Repeated in SeqNo=5
.:130 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
.:152 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
.:179 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
.:179 M:254 (E) No. of Bases conflict, LENGTH:Input:0 Counted:36 SEQ:8 ✓
.:207 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
.:207 M:254 (E) No. of Bases conflict, LENGTH:Input:0 Counted:51 SEQ:9 ✓
.:236 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
l:341 Repeated in SeqNo=10
.:264 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
l:341 Repeated in SeqNo=11
.:298 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
.:331 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
l:341 Repeated in SeqNo=15
.:346 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
l:341 Repeated in SeqNo=16
.:361 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
l:341 Repeated in SeqNo=17
.:381 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
.:381 M:254 (E) No. of Bases conflict, LENGTH:Input:0 Counted:22 SEQ:18 ✓
.:394 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
.:408 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
.:422 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
l:341 Repeated in SeqNo=21
.:437 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
l:341 Repeated in SeqNo=22
.:453 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
l:341 Repeated in SeqNo=23
.:470 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
l:341 Repeated in SeqNo=24
.:485 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
l:341 Repeated in SeqNo=25
.:500 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
l:341 Repeated in SeqNo=26
.:517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
l:341 Repeated in SeqNo=27
.:532 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
l:341 Repeated in SeqNo=28
.:547 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0
l:341 Repeated in SeqNo=29
.:563 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
l:341 Repeated in SeqNo=30
.:579 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0
l:341 Repeated in SeqNo=31
.:595 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0

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1:341 Repeated in SeqNo=32
 1:611 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0
 1:341 Repeated in SeqNo=33
 1:628 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0
 1:341 Repeated in SeqNo=34
 1:1069 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:23 SEQ:60 ✓
 1:1655 M:258 E: Mandatory Feature missing, <220> Tag not found for SEQ ID#:98 ✓
 1:1656 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:51 SEQ:98 ✓
 1:7201 M:252 E: No. of Seq. differs, <211> LENGTH:Input:30 Found:31 SEQ:379 ✓
 1:7417 M:254 E: No. of Bases conflict, LENGTH:Input:19 Counted:18 SEQ:388 ✓
 1:7417 M:252 E: No. of Seq. differs, <211> LENGTH:Input:19 Found:18 SEQ:388 ✓
 1:7523 M:254 E: No. of Bases conflict, LENGTH:Input:43 Counted:44 SEQ:393 ✓
 1:7523 M:252 E: No. of Seq. differs, <211> LENGTH:Input:43 Found:44 SEQ:393 ✓
 1:7587 M:254 E: No. of Bases conflict, LENGTH:Input:27 Counted:17 SEQ:396 ✓
 1:7587 M:252 E: No. of Seq. differs, <211> LENGTH:Input:27 Found:17 SEQ:396 ✓
 1:22 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (377) Counted (410)

Use of n and / or Xaa has been detected in the
 Sequence Listing. Review the Sequence Listing
 to ensure a corresponding explanation is present
 in the <220> to <223> fields of each sequence
 using n or Xaa.